

IN THE CLAIMS:

Please amend the claims by canceling Claims 1-9, 17, 20 and 74-79, without prejudice, as drawn to non-elected subject matter, canceling Claims 23-31 and 73, without prejudice, and adding new Claims 80-98 as follows:

1.-9. (Canceled)

17. (Canceled)

20. (Canceled)

23.-31. (Canceled)

73.-79. (Canceled)

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Contd. 80. (New) A method of assessing the effectiveness of a non nucleoside reverse transcriptase inhibitor ("NNRTI") on an HIV-1-infected patient, comprising: detecting, in a biological sample of the HIV-1-infected patient, the presence of a nucleic acid that exhibits a mutation at codon 230 of a nucleotide sequence encoding HIV-1 reverse transcriptase, wherein the presence of such a mutation correlates with a decrease in susceptibility to delavirdine or nevirapine.

2 81. (New) The method of claim 80, wherein the presence of the mutation at codon 230 correlates with a decrease in susceptibility to delavirdine and nevirapine.

3 82. (New) The method of claim 80, wherein the presence of the mutation at codon 230 correlates with a decrease in susceptibility to delavirdine.

4 83. (New) The method of claim 80, wherein the presence of the mutation at codon 230 correlates with a decrease in susceptibility to nevirapine.

5 84. (New) The method of claim 80, wherein the mutation at codon 230 encodes a leucine (L).

4_{85.} (New) The method of claim 80, further comprising evaluating whether the biological sample of the HIV-1-infected patient comprises a nucleic acid encoding HIV-1 reverse transcriptase having a mutation at codon 181.

7_{86.} (New) The method of claim 85, wherein the mutation at codon 181 encodes a cysteine (C).

8_{87.} (New) The method of claim 80, wherein the HIV-1-infected patient is being treated with an antiretroviral agent.

9_{88.} (New) A method of assessing the effectiveness of a non nucleoside reverse transcriptase inhibitor ("NNRTI") on an HIV-1-infected patient comprising detecting, in a biological sample of the HIV-1-infected patient, the presence of a mutation at codon 181 of the nucleic acid encoding HIV-1 reverse transcriptase, wherein the presence of such a mutation correlates with a decrease in susceptibility to delavirdine or nevirapine and little or no change in susceptibility to efavirenz.

10_{89.} (New) The method of claim 88, wherein the presence of the mutation at codon 181 correlates with a decrease in susceptibility to delavirdine and nevirapine and little or no change in susceptibility to efavirenz.

11_{90.} (New) The method of claim 88, wherein the presence of the mutation at codon 181 correlates with a decrease in susceptibility to delavirdine and little or no change in susceptibility to efavirenz.

12_{91.} (New) The method of claim 88, wherein the presence of the mutation at codon 181 correlates with a decrease in susceptibility to nevirapine and little or no change in susceptibility to efavirenz.

13_{92.} (New) The method of claim 88, wherein the mutation at codon 181 codes for a cysteine (C).

14_{93.} (New) The method of claim 88, further comprising evaluating whether the biological sample of the HIV-1-infected patient comprises nucleic acid encoding HIV-1 reverse transcriptase having a mutation at at least one of codon 98, codon 106 or codon 227.

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94. (New) The method of claim 93, wherein the mutation at codon 98 encodes a glycine (G).

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95. (New) The method of claim 93, wherein the mutation at codon 106 encodes an alanine (A).

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96. (New) The method of claim 93, wherein the mutation at codon 227 encodes a leucine (L).

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97. (New) The method of claim 93, wherein the mutation at codon 98 encodes a glycine (G); the mutation at codon 106 encodes an alanine (A); and the mutation at codon 227 encodes a leucine (L).

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98. (New) The method of claim 88 wherein the HIV-1-infected patient is being treated with an antiretroviral agent.